



1600

RAW SEQUENCE LISTING DATE: 09/12/2003
 PATENT APPLICATION: US/08/887,505A TIME: 14:25:45

Input Set : A:\47508-250(HYZ-040 CIP).ST25.txt
 Output Set: N:\CRF4\09122003\H887505A.raw

SEQUENCE LISTING

4 (1) GENERAL INFORMATION:
 6 (i) APPLICANT: Kilkuskie, Robert E.
 7 Frank, Bruce L.
 8 Goodchild, John
 9 Wolfe, Jia L.
 10 Roberts, Peter C.
 11 Hamlin, Jr., Henry A.
 12 Roberts, Noel A.
 13 Walther, Debra M.
 15 (ii) TITLE OF INVENTION: Oligonucleotides Specific for
 16 Hepatitis C Virus
 18 (iii) NUMBER OF SEQUENCES: 173
 20 (iv) CORRESPONDENCE ADDRESS:
 21 (A) ADDRESSEE: Hale and Dorr LLP
 22 (B) STREET: 60 State Street
 23 (C) CITY: Boston
 24 (D) STATE: MA
 25 (E) COUNTRY: USA
 26 (F) ZIP: 02109
 28 (v) COMPUTER READABLE FORM:
 29 (A) MEDIUM TYPE: Floppy disk
 30 (B) COMPUTER: IBM PC compatible
 31 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 32 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
 34 (vi) CURRENT APPLICATION DATA:
 C--> 35 (A) APPLICATION NUMBER: US/08/887,505A
 C--> 36 (B) FILING DATE: 02-Jul-1997
 37 (C) CLASSIFICATION:
 39 (vii) PRIOR APPLICATION DATA:
 40 (A) APPLICATION NUMBER: US 08/471,968
 41 (B) FILING DATE: 06-JUN-1995
 43 (viii) ATTORNEY/AGENT INFORMATION:
 44 (A) NAME: Kerner, Ann-Louise
 45 (B) REGISTRATION NUMBER: 33,523
 46 (C) REFERENCE/DOCKET NUMBER: 47508-250HYZ-040CIP
 48 (ix) TELECOMMUNICATION INFORMATION:
 49 (A) TELEPHONE: (617) 526-6000
 50 (B) TELEFAX: (617) 526-5000

**Does Not Comply
 Corrected Diskette Needed**

ERRORED SEQUENCES

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1800 (2) INFORMATION FOR SEQ ID NO: 98:
 1802 (i) SEQUENCE CHARACTERISTICS:
 1803 (A) LENGTH: 12 base pairs
 1804 (B) TYPE: nucleic acid
 1805 (C) STRANDEDNESS: single
 1806 (D) TOPOLOGY: linear
 W--> 1808 (ii) MOLECULE TYPE: DNA/RNA
 1810 (iii) HYPOTHETICAL: NO
 1812 (iv) ANTI-SENSE: YES
 1814 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 98:
 E--> 1816 UUCCGCAGAC CA

10/12

VERIFICATION SUMMARY

DATE: 09/12/2003

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Input Set : A:\47508-250(HYZ-040 CIP).ST25.txt

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L:35 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:36 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:62 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1
L:80 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2
L:98 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3
L:117 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4
L:134 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5
L:152 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6
L:170 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7
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L:206 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9
L:224 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10
L:242 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11
L:260 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12
L:278 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13
L:296 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14
L:314 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15
L:332 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16
L:350 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17
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L:440 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22
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L:494 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=25
L:512 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=26
L:530 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=27
L:548 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28
L:566 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29
L:584 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30
L:602 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31
L:620 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=32
L:638 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=33
L:656 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34
L:674 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=35
L:692 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36
L:710 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37
L:728 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38
L:746 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39
L:764 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=40
L:782 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=41
L:800 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=42
L:818 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=43
L:836 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=44
L:854 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=45
L:872 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=46

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L:908 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=48
L:926 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=49
L:944 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=50
L:1816 M:254 E: No. of Bases conflict, Input:10 Counted:12 SEQ:98